AMENDMENTS TO THE SPECIFICATION

Following the title, please replace the first paragraph with the following:

CROSS REFERENCE TO RELATED APPLICATIONS

This application is a Rule 371 national phase of International Application No. PCT/AU2004/001480, filed October 27, 2004, which was published in English as International Publication No. WO 2005/040198, and claims the benefit of Australian Application No. 2003905932, filed October 27, 2003. The entire disclosures of both applications are incorporated by reference herein.

Please amend the specification as shown:

Please delete the paragraph on page 6, line 4 and replace it with the following paragraph:

Figure 6 shows the amino acid sequence of βc (SEQ ID NO: 1).

Please delete the paragraphs on page 12, lines 23 to page 14, line 24 and replace them with the following paragraphs:

In another embodiment of the present invention, it is preferred that the motif is selected from any one of the following sequences:

NGPYLG.....PP..HSRSLP (SEQ ID NO: 2)

NVHYRT......P...KTHTMP (SEO ID NO: 3)

**RYFTQKEE.....TESG\$GP (SEQ ID NO: 4)

NKKYELODRDVCE....P.RYRSVSEP (SEO ID NO: 5)

NPTYSVM.....RSHSYP (SEQ ID NO: 6)

NIFYLIR...KSGSFPMPELKLSISFP (SEQ ID NO: 7)

NEEYLDLSQ......PLEQYSPSYP (SEQ ID NO: 8)

NQEYLDLSM......PLDQYSPSFP (SEQ ID NO: 9)

NATYKVD......VIQRTR\$KP (SEQ ID NO: 10)

NPEY.....HSASSGP (SEQ ID NO: 11)

NPDY.....WNHSLP (SEQ ID NO: 12)

NPSYSSNPFVNYN....KTSICSKSNP (SEO ID NO: 13)

NTLY.....FNSQSSP (SEQ ID NO: 14)

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NPVYQKTTEDEVHI...CHNQDGYSYP (SEQ ID NO: 15)
NPVYLKTTEEDLSIDIG..RH.SASVG (SEQ ID NO: 16)
NPTYKMYEGGEPDDVGGLLDADFALDPDKPTNFTNPVY (SEQ ID NO: 17)
NPIY.....KSAVTTVV (SEQ ID NO: 18)
NPLY.....KSAITTTV (SEQ ID NO: 19)
NPLY.....KEATSTFT (SEQ ID NO: 20)
NPLY.....RKPISTHT (SEQ ID NO: 21)
NPLY.....RGSTSTFK (SEQ ID NO: 22)
PGHYL.....RCDSTQP (SEQ ID NO: 23)
VOTYVLO.....GDPRAVSTOP (SEQ ID NO: 24)
QVLYGQLL.....GSPTSP (SEQ ID NO: 25)
HSGYRHQVPSVQVF.....SRSESTQP (SEQ ID NO: 26)
WKMYEVYDA.....KS.KSVSLP (SEQ ID NO: 27)
KIPYFHA.....GGS.KCSTWP (SEQ ID NO: 28)
ELDYCLKGLKL......P.S.RTWSPP (SEO ID NO: 29)
SGDYMPM.....SPKSVSAP (SEQ ID NO: 30)
SFYYSEENKLPEPEELDLEPENMESVP(LDPSASSSLP) (SEQ ID NO: 31)
EEIYIIM.....QSCWAFDSRKRPSFP (SEQ ID NO: 32)
ISQYLQN.....S.KRKSRP (SEQ ID NO: 33)
GTAY.....GLSRSQP (SEQ ID NO: 34)
***YLPQEDWAP......TSLTRP (SEQ ID NO: 35)
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LVAYIAFKRWNSCKQN...KQGANSRPVNQTPPPEGEKLHSDSGIS (SEQ ID NO: 36)

The sequences may correspond to the following:

betaR	NGPYLGPPHSRSLP (SEQ ID NO: 2)
Acetylcholine R	NVHYRTPKTH™MP (SEQ ID NO: 3)
Acetylcholine R alpha-5	**RYFTQKEETESGSGP (SEQ ID NO: 4)
C-C chemokine receptor 6	NKKYELQDRDVCEP.RYRSVSEP (SEQ ID NO: 5)
Middle T antigen	RPTYSVMRSHSYP (SEQ ID NO: 6)
integrin alpha 1	NIFYLIRKSGSFPMPELKLSISFP (SEQ ID NO: 7)
FGFR2 (KGF R)	NEEYLDLSQPLEQYSPSYP (SEQ ID NO: 8)
FGFR1 (flg)	NQEYLDLSMPLDQYSPSFP (SEQ ID NO: 9)
FGFR5	NATYKVDVIQRTRSKP (SEQ ID NO: 10)
Erb4	NPEYHSASSGP (SEQ ID NO: 11)
Erb4 (second)	%PDYWNHSLP (SEQ ID NO: 12)
Vaccinia virus protein A36R	NPSYSSNPFVNYNKTSICSKSNP (SEQ ID NO: 13)
Macrophage mannose R (MRC1)	NTLYFNSQSSP (SEQ ID NO: 14)
LDLR	NPVYQKTTEDEVHICHNQDGYSYP (SEQ ID NO: 15)
VLDL (rat)	NPVYLKTTEEDLSIDIGRH.SASVG (SEQ ID NO: 16)
LRP1 low density lipoprotein receptor-related protein 1 NPT	KMYEGGEPDDVGGLLDADFALDPDKPTNF\\NPVY (SEQ ID NO: 17)
integrin beta 1	NPIYKSAVT∜VV (SEQ ID NO: 18) (end of protein)
interin beta 7	NPLYKSAITTTV (SEQ ID NO: 19) (end of protein)
integrin beta 3	NPLYKEATSTFT (SEQ ID NO: 20) (end of protein)
integrin beta 5	RKPISTHT (SEQ ID NO: 21) (end of protein)
integrin beta 6	NPLYRGSTSTFK (SEQ ID NO: 22)

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G-CSFR1 (second) PGHYL.....RCDSTQP (SEQ ID NO: 23) G-CSFR1 VQTYVLQ.....GDPRAVSTQP (SEQ ID NO: 24) g-csf-r QVLYGQLL.....GSPTSP (SEQ ID NO: 25) IL-6B (gp130) HSGYRHQVPSVQVF.....SRSESTQP (SEQ ID NO: 26) WKMYEVYDA.....KS.KSVSLP (SEQ ID NO: 27) leptinR. KIPYFHA.....GGS.KCSTWP (SEQ ID NO: 28) prolactinR... ELDYCLKGLKL......P.S.RTWSPP (SEQ ID NO: 29) insulinR SGDYMPM.....SPKSVSAP (SEQ ID NO: 30) irs-1 SFYYSEENKLPEPEELDLEPENMESVP(LDPSASSSLP) (SEQ ID NO: 31) IGFI R EEIYIIM.....QSCWAFDSRKRPSFP (SEQ ID NO: 32) flt3 R

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 VEGFR2 (FLK1)
 ISQYLQN......S.KRKSRP (SEQ ID NO: 33)

 PDGF R-alpha
 GTAY.......GLSRSQP (SEQ ID NO: 34)

 IL-9R
 ****YLPQEDWAP......TSLTRP (SEQ ID NO: 35)

p75 NTR

LVAYIAFKRWNSCKQN...KQGANSRPVNQTPPPEGEKLHSDSGIS(phosphorylated)(SEQ ID NO: 36)

More particularly, the binding motif or an equivalent may have any one of the following sequences.

NGPY (**SEQ ID NO: 37**)

NVHY (SEQ ID NO: 38)

**RY

NKKY (SEQ ID NO: 39)

NPTY (SEQ ID NO: 40)

NIFY (**SEQ ID NO: 41**)

NEEY (SEQ ID NO: 42)

NQEY (SEQ ID NO: 43)

NATY (SEQ ID NO: 44)

NPEY (SEQ ID NO: 45)

NPDY (SEQ ID NO: 46)

NPSY (SEQ ID NO: 47)

NTLY (**SEQ ID NO: 48**)

NPVY (**SEQ ID NO: 49**)

NPIY (**SEQ ID NO: 50**)

NPLY (SEQ ID NO: 51)